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OM nucleic - nucleic search, using sw model

Run on: June 6, 2001, 20:03:40 ; Search time 182.07 Seconds
(without alignments)
7055.888 Million cell updates/sec

Title: US-09-494-297-1

Perfect score: 2274
Sequence: 1 atgaaaaaacaggttcc.....ggataagaacatgactag 2274

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 470324 seqs, 282468253 residues

Total number of hits satisfying chosen parameters: 940648

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

Pending Patents_NA_New:*
1: /cgnl_7/ptodata/2/pna/US06_NEW_COMB.seq:*
2: /cgnl_7/ptodata/2/pna/US07_NEW_COMB.seq:*
3: /cgnl_7/ptodata/2/pna/US08_NEW_COMB.seq:*
4: /cgnl_7/ptodata/2/pna/US09_NEW_COMB.seq:*
6: /cgnl_7/ptodata/2/pna/US60_NEW_COMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2274	100.0	2274	US-09-494-297-1	Sequence 1, Appl1
2	720.8	31.7	2229	US-09-494-297-3	Sequence 3, Appl1
3	51.2	2.3	902	US-09-739-449-1979	Sequence 1929, Ap
4	49.4	2.2	780	US-09-739-449-451	Sequence 451, Ap
5	48.6	2.1	69513	US-60-248-823-16	Sequence 16, Appl
6	47.8	2.1	753	US-09-739-449-1234	Sequence 1234, Ap
7	47.4	2.1	5897	US-08-956-171C-320	Sequence 320, App
8	47.2	2.1	1131	US-09-739-449-1953	Sequence 1953, Ap
9	46.4	2.0	723	US-09-739-449-8012	Sequence 8012, Ap
10	46.4	2.0	892	US-09-739-449-1335	Sequence 1335, Ap
11	46.4	2.0	2850	US-09-663-882-1	Sequence 1, Appl1
12	45.6	2.0	1250	US-09-739-449-1859	Sequence 1859, Ap
13	45.6	2.0	2841	US-08-956-171C-917	Sequence 177, App
14	44.8	2.0	1220	US-08-956-171C-178	Sequence 918, App
15	44.4	2.0	757	US-09-739-449-2286	Sequence 2286, Ap
16	44.4	2.0	1040	US-09-739-449-2287	Sequence 2287, Ap
17	44.4	2.0	787	US-09-739-449-2501	Sequence 2501, Ap
18	44.4	1.9	798	US-09-739-449-2641	Sequence 2641, Ap
19	44.4	1.9	786	US-09-739-449-1368	Sequence 1368, Ap
20	43.8	1.9	543	US-09-739-449-8029	Sequence 8029, Ap
21	43.6	1.9	856	US-09-833-343-74	Sequence 74, Appl
22	43.4	1.9	499	US-09-739-449-1332	Sequence 1332, Ap
23	43.4	1.9	776	US-09-739-449-2257	Sequence 2257, Ap
24	43.4	1.9	909	US-09-739-449-636	Sequence 636, App
25	43.4	1.9	117358	US-60-248-505-567	Sequence 567, App
26	42.8	1.9	1006	US-09-739-449-621	Sequence 621, App
27	42.8	1.9	1006	US-09-739-449-621	Sequence 621, App

28	42.8	1.9	43546	US-60-254-168-38	Sequence 38, Appl
29	42.6	1.9	780	US-09-739-449-778	Sequence 778, App
30	41.8	1.8	548	US-09-739-449-1457	Sequence 1457, Ap
31	41.8	1.8	403789	US-60-248-505-197	Sequence 197, App
32	41.6	1.8	681	US-09-739-449-2434	Sequence 2434, Ap
33	41.4	1.8	14333	US-08-956-171C-57	Sequence 57, Appl
34	41	1.8	385	US-09-737-223-6963	Sequence 6963, Ap
35	41	1.8	186647	US-60-248-823-22	Sequence 22, Appl
36	40.6	1.8	831	US-09-739-449-285	Sequence 285, App
37	40.6	1.8	1418	US-09-739-449-1613	Sequence 1613, Ap
38	40.2	1.8	664	US-09-739-449-739	Sequence 739, App
39	40	1.8	6373	US-08-956-171C-227	Sequence 227, App
40	39.8	1.8	6060	US-08-956-171C-534	Sequence 534, App
41	39.8	1.8	9821	US-08-956-171C-470	Sequence 470, App
42	39.8	1.8	15350	US-60-248-823-6	Sequence 6, Appl1
43	39.6	1.7	778	US-09-739-449-2435	Sequence 2435, App
44	39.6	1.7	1014	US-09-739-449-1697	Sequence 1697, Ap
45	39.4	1.7	849	US-09-739-449-1463	Sequence 1463, Ap

ALIGNMENTS

RESULT 1
US-09-494-297-1
Sequence 1, Application US/09494297
GENERAL INFORMATION:
APPLICANT: PODBIELSKI, ANDREAS
TITLE OF INVENTION: COLLAGEN-BINDING PROTEINS FROM STREPTOCOCCUS PYOGENES
FILE REFERENCE: P06628US0/BAS
CURRENT APPLICATION NUMBER: US/09/494,297
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1
LENGTH: 2274
TYPE: DNA
ORGANISM: Streptococcus pyogenes
US-09-494-297-1

Query Match	100.0%	Score 2274;	DB 5;	Length 2274;
Best Local Similarity	100.0%	Pred. No. 0;		
Matches 2274;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
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DB	1	atgaaaaaacaggttccaaataagcttaacttaatactcaaaaggtatgaagt	60	
QY	61	aaaaactcaaacagattactgtcaacttaagtggaggtcttttaagtctcgcttg	120	
DB	61	aaaaactcaaacagattactgtcaacttaagtggaggtcttttaagtctcgcttg	120	
QY	121	gtacttccatgttggtgtaagactgttttggttaagataatccctcgaagcaaac	180	
DB	121	gtacttccatgttggtgtaagactgttttggttaagataatccctcgaagcaaac	180	
QY	181	gcataataatccagatcgaatcgaatcgaatcgaatcgaatcgaatcgaatcgaat	240	
DB	181	gcataataatccagatcgaatcgaatcgaatcgaatcgaatcgaatcgaatcgaat	240	
QY	241	gggcatccatataataaacagtttagagtagcacacagattaaagggttaactgaaga	300	
DB	241	gggcatccatataataaacagtttagagtagcacacagattaaagggttaactgaaga	300	
QY	301	agtagaagttataaagtttagtttaatttaagaagaacattcctcgcgaataagat	360	
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QY	361	agtaggtttaaaaaagtgtaaaaaaacatgagatctctcaaaaatttgaagaatt	420	
DB	361	agtaggtttaaaaaagtgtaaaaaaacatgagatctctcaaaaatttgaagaatt	420	

QY	421	ggagtagcccttagaattacgagggagatgagctaaatcaagaagtacagctgttatg	480
Db	421	ggatggagcccttaaatatcacgggagatgagctaaatcaagaagtacagctgttatg	480
QY	481	aatggacatccacaanaatgccaatgagctatatggaagcctgtggaacccctggaatgc	540
Db	481	aatggacatccacaanaatgccaatgagctatatggaagcctgtggaacccctggaatgc	540
QY	541	agaagtacaagaagcggtatgtgttactatctgtataatgtctcatlttcaatccaag	600
Db	541	agaagtacaagaagcggtatgtgttactatctgtataatgtctcatlttcaatccaag	600
QY	601	gaagaatttaaaagggagtcagaagaatgtaactgtgtatgtactctctaattctttgag	660
Db	601	gaagaatttttaaagggagtcagaagaatgtaactgtgtatgtactctctaattctttgag	660
QY	661	cgtaagctttggaagcaactatgtaaccgaatttggcaactaaatgccaacaagaat	720
Db	661	cgtaagctttggaagcaactatgtaaccgaatttggcaactaaatgccaacaagaat	720
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Db	721	ccggaatgtttcagactaagatatttttgaagctctgaggaacaaggagataataa	780
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Db	781	ggatacccaaatcttttgaagtggtgtgtgttgcctctactaaaccaactccagagac	840
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Db	841	ccaccaatgagctccaatgcaaccccaacgaagcctcagtaactataaagaatgtcata	900
QY	901	gggtatctctctaaattgctgtgaagtgccaacatlaaagttgacagggataacgtga	960
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Db	961	agtttcaagcgagagtggttttaagcagtaatgatatgtggaagaagaattgaactaca	1020
QY	1021	ggaacttacttaactaactgtaattatctccagctgtgtatgatacgagagccaatc	1080
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QY	1081	acttttaaggttgaagcctgagcaagtgatatacttlatgtatggaanaacagattga	1140
Db	1081	acttttaaggttgaagcctgagcaagtgatatacttlatgtatggaanaacagattga	1140
QY	1141	cccaataaaggatagtgtagagccttactccagtagagacataatgttttgaagaatt	1200
Db	1141	cccaataaaggatagtgtagagccttactccagtagagacataatgttttgaagaatt	1200
QY	1201	agcggtttaactacaacaactatgcaaaattttatatagcgaanaataaataatgga	1260
Db	1201	agcggtttaactacaacaactatgcaaaattttatatagcgaanaataaataatgga	1260
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Db	1261	tcacaggttgtctatgtctttaaagtcagatcaaaatctccacogaagcttgaagat	1320
QY	1321	gggaaaaacaatgactccagacttttacacaaggaagtaaaatacacatcatgtcag	1380
Db	1321	gggaaaaacaatgactccagacttttacacaaggaagtaaaatacacatcatgtcag	1380
QY	1381	cgtagccctttaaatactgctgaaaccaagagataccgataccgtaaccttcttaaa	1440
Db	1381	cgtagccctttaaatactgctgaaaccaagagataccgataccgtaaccttcttaaa	1440
QY	1441	catatcaaaaaagtaattggaagaaggtttacacggggaanaagacacagctattgata	1500
Db	1441	catatcaaaaaagtaattggaagaaggtttacacggggaanaagacacagctattgata	1500
QY	1501	ggtctaacctggaacaatgycgtgcggtactccagttgaagaatataatttcaactga	1560

Dh	1501	ggtctaacctgagacacaatttgcgtggtgcgtactcgttttagccaattatatttccaacgt	1560
Qy	1561	agtgctgaattagaataagataaactaaagaactacatcgtgttttggagacatgatat	1620
Dh	1561	agtgctgaattagaataagataaactaaagaactacatcgtgttttggagacatgatat	1620
Qy	1621	agtaacttaagcagtttgcataaaaccccttgcataatcgcctcaagataagtaaccccccag	1680
Dh	1621	agtaacttaagcagtttgcataaaaccccttgcataatcgcctcaagataagtaaccccccag	1680
Qy	1681	ctaacctgacctgatatcttccttaccgaaataacaataacatacctcttatcgtgaact	1740
Dh	1681	ctaacctgacctgatatcttccttaccgaaataacaataacatacctcttatcgtgaact	1740
Qy	1741	caatgtagcaccgcagaagaatttagttgatatatttcgtatggaagataaaaaagaagtata	1800
Dh	1741	caatgtagcaccgcagaagaatttagttgatatatttcgtatggaagataaaaaagaagtata	1800
Qy	1801	ccgttaactcctaatttaataaatgtagaanaaacggtgacgtgttgcgtgcgtgagacaact	1860
Dh	1801	ccgttaactcctaatttaataaatgtagaanaaacggtgacgtgttgcgtgcgtgagacaact	1860
Qy	1861	aaagatttccattttggaatttgaattaaataaataaataagcaagaatttgccttccaact	1920
Dh	1861	aaagatttccattttggaatttgaattaaataaataaataagcaagaatttgccttccaact	1920
Qy	1921	gttaaaacagataaaaacaaaccccgaaatttaagaatgtraaagcacaacttaattttaa	1980
Dh	1921	gttaaaacagataaaaacaaaccccgaaatttaagaatgtraaagcacaacttaattttaa	1980
Qy	1981	catggggaagaagtttaacaactccaagtttataccagaaggttatcttccataccttgcacaaga	2040
Dh	1981	catggggaagaagtttaacaactccaagtttataccagaaggttatcttccataccttgcacaaga	2040
Qy	2041	acagatctctgaaggcctataaaggttaaaagttaataagccaaagaagttagcacaatgctacagtt	2100
Dh	2041	acagatctctgaaggcctataaaggttaaaagttaataagccaaagaagttagcacaatgctacagtt	2100
Qy	2101	tcaaaaacagaataaacaanaatgtagtgagacactgctcttggaaataataaagaagccgtt	2160
Dh	2101	tcaaaaacagaataaacaanaatgtagtgagacactgctcttggaaataataaagaagccgtt	2160
Qy	2161	gttccctacagagagttgataccaagaatccaatggtcgtatcctatgtagtatgcgtgt	2220
Dh	2161	gttccctacagagagttgataccaagaatccaatggtcgtatcctatgtagtatgcgtgt	2220
Qy	2221	atcaggtttgggagatctggggaattcaacacagataagaataagaanaacatgactag	2274
Dh	2221	atcaggtttgggagatctggggaattcaacacagataagaataagaanaacatgactag	2274

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RESULT 2
US-09-494-297-3
; Sequence 3, Application US/09494297
; GENERAL INFORMATION:
; APPLICANT: PODBIELSKI, ANDREAS
; TITLE OF INVENTION: COLLAGEN-BINDING PROTEINS FROM STREPTOCOCCUS PYOGENES
; FILE REFERENCE: P06628050/BAS
; CURRENT APPLICATION NUMBER: US/09/494,297
; CURRENT FILING DATE: 2000-01-31
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 3
; LENGTH: 2229
; TYPE: DNA
; ORGANISM: Streptococcus pyogenes
US-09-494-297-3

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Query Match	31.7%;	Score 720.8;	DB 5;	Length 2229;
Best Local Similarity	61.6%;	Pred. No. 3.3e-140;		
Matches 1372; Conservative	0;	Mismatches 787;	Indels 69;	Gaps 11;

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DB 40 aacaaagacgacaaacagcagtcgacttgaagatatttgacgtttgtagcttg 99
QY 121 gtaactccatgtgtgtgtaagacgtcttttggtttagtagaatctcagccaaac 180
DB 100 ataggaatagtaggtttcttcaacagagcgttcggagctgaagaacaaatcagtaacaaa- 158
QY 181 gcaataaatccagatcaagtctcggaatacagatgtaatgataatgataatgataa 240
DB 159 -----tagacaagctcaattcaagatcacgttgcgtatgctatgattcttaccataa 213
QY 241 gggcatccatataataaacggtttagtagtgcacagatctaagggttaacttagaaga 300
DB 214 ggtaccacagactatagtccgtttaaagaacttaccataatttaaaagttaattagaaga 273
QY 301 agtaagaattacagatttcttgaatttaagaagaagcattctctcgcgagat 360
DB 274 agtaagaattacagatctcgtcttaatttaacaaacatttccatccaagtcagat 333
QY 361 agtagtgttaaaaagtgtataaaaacatgaltggaatctctcaaaaatttgaagattat 420
DB 334 agtgttagatcacatggtataaaaacttggaaggaactaatgaaactttatcaagttia 393
QY 421 ggcgtatgacctgaattacagggagatgagctaatcagaagttagagctgtatgtat 480
DB 394 gcaatataaaccaagaatagagaagcgagcttacaacaaatataatgagatctctat 453
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DB 454 aatggatatacttaataatcgtatggtatgataatgaagggatagatctcttaaaagctat 513
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DB 514 ttatgctcacaagtatgtattgtta-----taactgattccagctcaaatlaatccgat 567
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QY 1195 gaatttagcgttttaactacaacaaacatgccaacatttataatgcaaaaataaaat 1254
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QY 1600 gttttgagacatgaatgatagtactttagcaggttgcataaaatccttgaataagct 1659
DB 1555 ggaattgaatcattagtgataaaacccctagctgtccacaaagaatlaactatagct 1614
QY 1660 caagatagtatctccacagcttaacagacttgatttcttctcgaataacaataa 1719
DB 1615 caaaatgtagcgtccctcaactcaacaaatcttgatttcttgcglaactaataatagcaa 1674
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DB 1675 gacaaactcttatttggagaagaatgacacccaagatattgttgcgtatcgat 1734
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DB 1735 gaagataaaaagaagaagttattccagttacacaggttgcagtgagaaaaaaacagta 1794
QY 1837 actggtttagctgtgacgaactaaagatttccattttgaattgaataaaaaataat 1896
DB 1795 gtcggtgagttgggagataaaactaaaggttccaaatttgaacttgaattgaaagataa 1854
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QY 2017 ggttattcttaacttgcgaagaacagatctcgaagcctataagtttaagttatagc 2076
DB 1975 ggaatattcttatactcgaagaagcgtggaagctlaagattatagtaaacggttgaac 2034
QY 2077 caagaagtagcaaatgttacaagtttcaaaaacggaataacaggtgtatgtagcaactgtc 2136
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QY 2137 ttgaaaaataaagaagcgtgtgttctcagaggtgtatcnaaagatcaatgtgctat 2196
DB 2095 ttgtgaaaaccgaaaagatctgtccccaacaggttgttgcaacagatlggggtatctat 2154
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QY 2001 tcaaggttaccagaaggttatcttacctgttccaagaacagattctgaagctataa 2060
Db 483 AAAAAAAAAAAATNNNTAAAAAANNAAAAAANNAAAAAANAAAAAANAAAAA 424
QY 2061 gatttaagttatagccaagaagtagcaaatgtctacagttccaataaacagataacaag 2120
Db 423 AATTAAANAAAAAANAAAAAATTTAAAAAANAAAAAANAAAAAANNNTNANNNANNN 364
QY 2121 tgaatgacactgtctttgaaataataagaagcgtgtgtctcacaagagttgata 2180
Db 363 NNNNNNNAAAAAANAAAAAATTAATANTANNNTNNNNNNNNAAAAAANNNNNNNNNN 304
QY 2181 aaagatcaatgctata 2199
Db 303 NNNNNNNNNNTTCTCTA 285

RESULT 7
US-09-739-449-2256/c
; Sequence 2256, Application US/09739449
; GENERAL INFORMATION:
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
; FILE REFERENCE: 38-10(15490)C
; CURRENT APPLICATION NUMBER: US/09/739,449
; PRIOR FILING DATE: 2000-12-19
; PRIOR APPLICATION NUMBER: US 09/514,000
; PRIOR FILING DATE: 2000-02-23
; NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 2256
; LENGTH: 834
; TYPE: DNA
; ORGANISM: Agrobacterium tumefaciens
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(834)
; OTHER INFORMATION: unsure at all n locations
US-09-739-449-2256

Query Match 2.1%; Score 47.8; DB 5; Length 834;
Best Local Similarity 43.9%; Pred. No. 0.34;
Matches 196; Conservative 0; Mismatches 250; Indels 0; Gaps 0;

QY 1708 aatacaataataatcaatctcttattggaactcagtgycatccgaagaatttagat 1767
Db 660 AATTAATAAAAAAAAAATTAATATAAAAAAAAAAAAAAAAAAATAAATAAATAAATAAATAA 601
QY 1768 attatcgtatggaagataaagaagttataactgttaaccataatctaacttgga 1827
Db 600 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAATTAATAAAAAAAAAAAAAAAAAAAAAA 541
QY 1828 aaacagcgtactggttagctgtgacgaagaatctccatttggaaatgaatta 1887
Db 540 AAAAAAAAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAA 481
QY 1888 aaaaataataagcaagaattgcttctcaaacgtttaaacagataaacaacactgaa 1947
Db 480 AAAAAAAAAAAAAAAAAAATTAATAAAAAAAAAAAAAAAAAAATAAATAAATAAATAAATAA 421
QY 1948 tttaagaatggtgaagcaaccataatttaaacatggggaaagttaacctcaaggt 2007
Db 420 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAATAAATAAATAAATAAATAAATAAATAA 361
QY 2008 ttaccagaagttacttcttacctgtccaagaacagattctgaagctataagttaa 2067
Db 360 AAAAAAAAAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAA 301
QY 2068 gttaatagccaagaagtcgaatgtctcaaaacaggaatacaagtcagtga 2127
Db 300 AAAAAAAAAAAAAAAAAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAA 241

```

OY 2128 acacttgccttgaataataaga 2153
 | | | | |
 Db 240 CCGCCCCCCCCCAAAAAAAAAA 215

RESULT 8

US-08-956-171C-320
 ; Sequence 320, Application US/08956171C
 ; GENERAL INFORMATION:

APPLICANT: Charles Kunsch

Gil H. Choi

Patrick S. Dillon

Craig A. Rosen

Steven C. Barash

Michael R. Fannon

TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
 NUMBER OF SEQUENCES: 5255

CORRESPONDENCE ADDRESS:

ADDRESSEE: Human Genome Sciences, Inc.

STREET: 9410 Key West Avenue

CITY: Rockville

STATE: Maryland

COUNTRY: USA

ZIP: 20850

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 Inch, 1.4MB storage

COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

SOFTWARE: ASCII Text

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/956,171C

FILING DATE: 20-Oct-1997

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/009,861

FILING DATE: January 5, 1996

APPLICATION NUMBER: 08/781,986

FILING DATE: January 3, 1997

ATTORNEY/AGENT INFORMATION:

NAME: Hoover, Kenley K.

REGISTRATION NUMBER: 40,302

REFERENCE/DOCKET NUMBER: PB248P1

TELECOMMUNICATION INFORMATION:

TELEPHONE: (301) 610-5790

TELEFAX: (301) 309-8439

INFORMATION FOR SEQ ID NO: 320:

SEQUENCE CHARACTERISTICS:

LENGTH: 5897 base pairs

TYPE: nucleic acid

STRANDEDNESS: double

TOPOLOGY: linear

SEQUENCE DESCRIPTION: SEQ ID NO: 320:

US-08-956-171C-320

Query Match 2.1%; Score 47.4; DB 4; Length 5897;
 Best Local Similarity 46.3%; Pred. No. 0.6; Indels 0; Gaps 0;

Matches 156; Conservative 0; Mismatches 181; Indels 0; Gaps 0;

OY 865 caaacactcagctactttagaagatgtctatagtgatgattactctaattgtctgaa 924
 | | | | |

Db 4351 CCAATATATTCGATTAGATTGTAATCAAAAAGAAAGTAATCAATTTTAGATTATT 4410
 | | | | |

OY 925 ggtgcacactcagctgcaggggataacgtgatatgttttcaagcgaagtgattacg 984
 | | | | |

Db 4411 GATGAATCATTTTCGTAAGATGATGAACTTTTATCAAGAGTCTCTCAATTA 4470
 | | | | |

OY 985 agtaatgatattggagaagaatgtaactcagatgtaagtaacttaactgaatg 1044
 | | | | |

Db 4471 ATTAAATGAATAAAGAAAGAGCCATGATGTTTATTTTAAACCTGATGATATT 4530
 | | | | |

OY 1045 aattccacgctggttatgtatcgcagagccatcatttaaggttgaaagtgcaaa 1104
 | | | | |

Db 4531 CCTAATACAGTTGGTAAGCGCATGAGAGACTAGCTAATTAAGTGTGATTAAT 4590
 | | | | |
 OY 1105 ggtatatactattatgtgaaacagattgaaatcccaataaagagatagtagccct 1164
 | | | | |
 Db 4591 GTTCATGCTGCTGCTGCGCTAAATAATGATGCTGAGCCATTAAAGATTAAAGAAACAT 4650
 | | | | |
 OY 1165 tactcagtagaagacataatgatttgaagaattta 1201
 | | | | |
 Db 4651 AATCAAGATACAAAATTAATTTCAGTAAACACAGCTTA 4687

RESULT 9

US-09-739-449-1953
 ; Sequence 1953, Application US/09739449

GENERAL INFORMATION:

APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof

FILE REFERENCE: 38-10(15490)C

CURRENT APPLICATION NUMBER: US/09/739,449

CURRENT FILING DATE: 2000-12-19

PRIOR APPLICATION NUMBER: US 09/514,000

PRIOR FILING DATE: 2000-02-23

NUMBER OF SEQ ID NOS: 13351

SEQ ID NO 1953

LENGTH: 1131

TYPE: DNA

ORGANISM: Agrobacterium tumefaciens

FEATURE:

NAME/KEY: unsure

LOCATION: (1)..(1131)

OTHER INFORMATION: unsure at all n locations

US-09-739-449-1953

Query Match 2.1%; Score 47.2; DB 5; Length 1131;
 Best Local Similarity 44.1%; Pred. No. 0.48;

Matches 163; Conservative 0; Mismatches 207; Indels 0; Gaps 0;

OY 1781 aagataaaaaagaagttatacctgtatactcataatttaacattgagaaaaagtgactg 1840
 | | | | |

Db 510 aaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaa 569
 | | | | |

OY 1841 gtttagctggtgcagacttaagatttcatttgaattgaattgaattgaattgaattga 1900
 | | | | |

Db 570 maaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaa 629
 | | | | |

OY 1901 aagaattgcttctcaactgtttaaacagataaaacaaactcgaaatttaagaatgta 1960
 | | | | |

Db 630 aaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaa 689
 | | | | |

OY 1961 aagcaaccataatttaaacatggtgaaagttaacacttaaggtttaccagaaggt 2020
 | | | | |

Db 690 aaaaaaagaanaaaaaaagaanaaaaaaagaanaaaaaaagaanaaaaaaagaana 749
 | | | | |

OY 2021 attcttacttgcagaagaacagattctgaaggtcataagtttaagtttaagtttaag 2080
 | | | | |

Db 750 aaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaaanaaaaaa 809
 | | | | |

OY 2081 aagtagcaatgctacagttcaaaaacaggaataaacaagtatgagacagctgtttg 2140
 | | | | |

Db 810 aagaaagcaanaagagagaaaaaagaagaagaagaagaagaagaagaagaagaaga 869
 | | | | |

OY 2141 aaaaataaa 2150
 | | | | |

Db 870 aaaaaaa 879

RESULT 10

US-09-739-449-8012
 ; Sequence 8012, Application US/09739449

GENERAL INFORMATION:

APPLICANT: Hinkle, Gregory J.

```

RESULT 12
US-09-663-882-1
; Sequence 1, Application US/09663882
; GENERAL INFORMATION:
; APPLICANT: MacInga, David R.
; TITLE OF INVENTION: Counter-selection vector for gram positive bacteria
; FILE REFERENCE: ID01008 US
; CURRENT APPLICATION NUMBER: US/09/663,882
; CURRENT FILING DATE: 2001-09-15
; PRIOR APPLICATION NUMBER: US 60/154,694
; PRIOR FILING DATE: 1999-09-17
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: PatentIn Ver. 2.1

```



```
; SEO ID NO 1
; LENGTH: 2850
; TYPE: DNA
; ORGANISM: Staphylococcus aureus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1355)..(2047)
; NAME/KEY: RBS
; LOCATION: (1344)..(1349)
; NAME/KEY: -10_signal
; LOCATION: (1302)..(1307)
; NAME/KEY: -35_signal
; LOCATION: (1281)..(1284)
; US-09-663-882-1
```

```
Query Match 2.0%; Score 46; DB 5; Length 2850;
Best Local Similarity 47.0%; Pred. No. 1;
Matches 142; Conservative 0; Mismatches 160; Indels 0; Gaps 0;
```

```
QY 865 caaacgactcagctactattagaagaatgctatagtgattactctaattgctgaa 924
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1370 ccaattctcattagatttgcattcaaaagaataaattcaattttagatttatt 1429
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 925 ggtgcacattacagttgacaggggatacgtgatatgttttcaagcgagagttagc 984
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1430 gataaatcattatcgtaaaagtgatgtaagtaactttttatcagaaggtcccaatta 1489
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 985 agtaatgatatctggagaagaattgaaactcagatggaacttaacttaactgaattg 1044
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1490 attaatgataaagaagaagagcgcatgattttagatttaaaacgtcatgatalct 1549
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1045 aattctcagctggttatagatcgacgagcgaactctttaaggttgtagctggcaaa 1104
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1550 cctaatacagctggtgaagcgatggaagcactgaattgactgttgtaaat 1609
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1105 gctatatacttattatgtaaaacagatggaatacccaataaagagatagtaagcct 1164
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1610 gtcatgctgctggtggtgcgaataatgctgtcgtgagccattaaagattaaagaacat 1669
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1165 ta 1166
DB 1670 aa 1671
```

```
RESULT 13
US-09-739-449-1859/C
; Sequence 1859, Application US/09739449
; GENERAL INFORMATION:
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
; FILE REFERENCE: 38-10(15490)C
; CURRENT APPLICATION NUMBER: US/09/739,449
; PRIOR FILING DATE: 2000-12-19
; PRIOR APPLICATION NUMBER: US 09/514,000
; PRIOR FILING DATE: 2000-02-23
; NUMBER OF SEQ ID NOS: 13351
; SEQ ID NO 1859
; LENGTH: 1250
; TYPE: DNA
; ORGANISM: Agrobacterium tumefaciens
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(1250)
; OTHER INFORMATION: unsure at all n locations
; US-09-739-449-1859
```

```
Query Match 2.0%; Score 45.6; DB 5; Length 1250;
Best Local Similarity 33.4%; Pred. No. 1;
Matches 204; Conservative 0; Mismatches 407; Indels 0; Gaps 0;
```

```
QY 1543 atatatattcactgtagtgctgaattagatagaataactaaagaactcatcgtgt 1602
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1181 AAAAAAAAAAANNTAANNNNNNANNAANNTAAAAAATTAANAAAAAANAAAAAANNNNA 1122
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1603 ttggagacatgagtagtactttagcagctgtcctaatactctgtagaatcgctcaa 1662
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1121 AAAAAAAAAAATTAANAAAAAANAAAAAANNNNNNNNNAANNAANNAANNNNNAAAAA 1062
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1663 gatagtatccctccacgctcaactgactgttctttatccgataacataatat 1722
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1061 NNNNAATTAANNAANAAAAAANNNNNNNNNAANNAANNAANNAANNAANNNN 1002
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1723 caatctcttatgtgaactcagtgcatccagaagatttagtgatatctcgtatgaa 1782
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 1001 AAAAAAAAAAANAAAAAANAAAAAANNNNAANNAANNNNNNNAAAAAANAAAAA 942
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1783 gataaaaaagaagttatcctgtactcaataattaacattgagaanaaacggtgactgt 1842
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 941 AAAAAAAAAAANNNNNNNAANNAANNAANNAANNAANNAANNAANNAANNAANNA 882
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1843 ttgctggtgacagactaaagattccatttgaattgaattgaataaataataagcaa 1902
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 881 ANAAAAAANNAANNAANAAAAAANNAANNAANNAANNAANNAANNAANNAANNA 822
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1903 gaattgcttctcaactgttaaacagatacaacaacctcgatttaagaatgtgtaa 1962
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 821 AANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNA 762
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 1963 gcaaccataatttaacaacatgggaaagttaacacttaaggtttaccagaagttat 2022
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 761 AAAAAAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNA 702
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 2023 tcttactgttcagaagaacagattctgaaggtataaggttaagttatagccaaga 2082
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 701 NNNNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNAANNA 642
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 2083 gtagcaatgctcagtttcaaaacagagaatacaagaatgtagacactgtcttgaa 2142
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 641 AAAAAAAAAAANAAAAAANAAAAAANAAAAAANAAAAAANAAAAAANNAANNA 582
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
QY 2143 aataataaga 2153
DB 581 AAAAAAAAAA 571
```

```
RESULT 14
US-08-956-171C-177
; Sequence 177, Application US/08956171C
; GENERAL INFORMATION:
; APPLICANT: Charles Kunsch
; APPLICANT: Gil H. Chol
; APPLICANT: Patrick S. Dillon
; APPLICANT: Craig A. Rosen
; APPLICANT: Steven C. Barash
; APPLICANT: Michael R. Fannon
; TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
; NUMBER OF SEQUENCES: 5255
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Human Genome Sciences, Inc.
; STREET: 9410 Key West Avenue
; CITY: Rockville
; STATE: Maryland
; COUNTRY: USA
; ZIP: 20850
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.50 inch, 1.4MB storage
; COMPUTER: HP Vectra 486/33
; OPERATING SYSTEM: MSDOS version 6.2
; SOFTWARE: ASCII Text
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/956,171C
; FILING DATE: 20-Oct-1997
; CLASSIFICATION: <Unknown>
```



```

; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/009,861
; FILING DATE: January 5, 1996
; APPLICATION NUMBER: 08/781,986
; FILING DATE: January 3, 1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Hoover, Kenley K.
; REGISTRATION NUMBER: 40,302
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (301) 610-5790
; TELEFAX: (301) 309-8439
; INFORMATION FOR SEQ ID NO: 177:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2841 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; SEQUENCE DESCRIPTION: SEQ ID NO: 177:
US-08-956-171C-177

Query Match          2.0%; Score 45.4; DB 4; Length 2841;
Best Local Similarity 51.2%; Pred. No. 1.3;
Matches 106; Conservative 0; Mismatches 101; Indels 0; Gaps 0;

QY 1710 taacataaatcaatcttattggaactcagtgcatccagaagaattagttgat 1769
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 689 TAAATAAGCATTAATCGTAATGTTAAATAAATGATTTAGAGCAAAACATAGATAT 748

QY 1770 tatcgtatggaagaataaagaagattataccgttaactatatttaaatggagaa 1829
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 749 TAAATGAAGTAAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 808

QY 1830 aacggtgactgttgaactggtgacagaactaaagattccatttgaattgaataa 1889
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 809 AGAAGGATTCATGATTAAGTAATCAAAATTCGAGATTGTCTTGTGGTGAGAGTACA 868

QY 1890 aataataagcaagaattgcttctca 1916
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 869 AAATCAGATATGACTTATGTTCTAA 895

RESULT 15
US-08-956-171C-918/c
; Sequence 918, Application US/08956171C
; GENERAL INFORMATION:
; APPLICANT: Charles Kunsch
;           Gil H. Choi
;           Patrick S. Dillon
;           Craig A. Rosen
;           Steven C. Barash
;           Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5255
CORRESPONDENCE ADDRESS:
; ADDRESSEE: Human Genome Sciences, Inc.
; STREET: 9410 Key West Avenue
; CITY: Rockville
; STATE: Maryland
; COUNTRY: USA
; ZIP: 20850
COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.50 inch, 1.4mb storage
; COMPUTER: HP Vectra 486/33
; OPERATING SYSTEM: MSDOS version 6.2
; SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/956,171C
; FILING DATE: 20-Oct-1997
; CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/009,861
```

```

; FILING DATE: January 5, 1996
; APPLICATION NUMBER: 08/781,986
; FILING DATE: January 3, 1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Hoover, Kenley K.
; REGISTRATION NUMBER: 40,302
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (301) 610-5790
; TELEFAX: (301) 309-8439
; INFORMATION FOR SEQ ID NO: 918:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1220 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; SEQUENCE DESCRIPTION: SEQ ID NO: 918:
US-08-956-171C-918

Query Match          2.0%; Score 44.8; DB 4; Length 1220;
Best Local Similarity 46.9%; Pred. No. 1.5;
Matches 136; Conservative 1; Mismatches 153; Indels 0; Gaps 0;

QY 1872 tttagaattgaattaaataataagcaagaattgcttcttaactglttaaacaga 1931
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 937 TTTAGAAATTCCACCAATTGTTGTACAAAATCAATTACTCAGAAAGTTAAACAGCGA 878

QY 1932 taacaaacactggaattaaagatggttaagcaaccatttaattaaacatgggaaag 1991
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 877 CATGATATTCGTATGATTAATGATGATGATGATGATGATGATGATGATGATGAT 818

QY 1992 tttaacacttaaggtttaccagaaggttattcttacctgtgcaagaagaattctga 2051
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 817 AATGCTTATCAAGATTAACGTCGCTTATTGCTGACAAAGAAAGATTAACAAAAC 758

QY 2052 aggtctataaggttaagtttaataagcaagaagtagcaaatgctcagtttcaaaacag 2111
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 757 ACGTATGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 698

QY 2112 aatacaagtgatgagacactgcttgaataataaagaagcctgtg 2161
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
DB 697 TACACCTATATGATTGCCAGGTGTTATTGAATAATGTCACAAAGTATCG 648
```

Search completed: June 6, 2001, 22:11:21
Job time: 7661 sec

